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(54) Title: RECOMBINANT IGF BINDING PROTEIN (IBP-1)

(I)

(57) Abstract

An IGF binding protein which has the amino acid sequence (I), or an equivalent modification thereof, such as a gly-cosylated modification. Further is indicated a DNA-sequence, coding for the protein, an expression vector and a pharmaceutical preparation containing the protein. The protein is effective as a potentiator for the function of IGF-compounds.

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Recombinant IGF binding protein (IBP-1)

The present invention relates to an Insulin-like Growth Factor Binding Protein, IBP-1, having a molecular weight of about 28 kD, derived from human placenta/endometrium, and equivalent modifications thereof.

The invention further relates to a DNA-structure, coding for the IBP-1, expression vectors containing this DNA-structure and procaryotic or eucaryotic cells comprising such a vector.

The invention still further relates to pharmaceutical preparations comprising IBP-1.

IGF or Insulin like Growth Factor is synonymous with somatomedins. The family of somatomedins are members of a group of polypeptides derived from the insulin 15 gene. The gene products include insulin, insuline-like growth factor (IGF)I and II, relaxin and the B-unit of nerve growth factor (NGF) (Rinderknecht and Humbel, 1978a; Rinderknecht and Humbel, 1978b; Bradshaw 1978; Isaacs et al., 1978). IGF elicits classical insulin 20 effects on all target tissues of insulin, i.e. IGF increases glucose metabolism of adipose tissue and stimulates lipid, glycogen, and protein synthesis. IGF also stimulates DNA synthesis in certain cell types. 25 This feature reflects the capacity of IGFs to induce cell proliferation and promote organ growth in vivo. Furthermore, IGF acts on differentiation of mesenchymal cells (Froesch et al., 1985).

IGF-I and IGF-II are unique in that they are complexed to specific binding proteins in plasma (Smith, 1984).

At least two different binding proteins have been iden-

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tified in adult human serum, namely (1) binding protein 53 (BP-53) a GH dependent binding protein, believed to be derived from the 150 kD complex which carries most of the endogenous IGF peptides, (2) IBP-1, an IGF binding protein of about 30-40 kD which is tissue specifically expressed in endometrium and liver and accounts for most of the unsaturable binding sites in plasma. While the 53 kD-binding protein is under GH control the 30-40 kD species appears to be expressed in a GH independent way.

The lower molecular weight binding protein was initially identified in human amniotic fluid and has been purified and characterized (Chochinov et al., 1977; Drop et al., 1979; Drop et al., 1982). This 30-40 kD IGF binding protein appears to be identical to binding proteins that have been purified from human serum and the human hepatoma cell line, HEPG2 (Drop et al., 1984a; Povoa et al., 1984; Povoa et al., 1985). Povoa et al showed that the NH₂-terminal amino acid sequence of the binding protein found in amniotic fluid and from the HEPG2 cell line are similar (Povoa et al., 1985).

Placental protein PP12, a protein originally isolated from human placenta, was found to bind IGF as well as to have an identical NH₂-terminal amino acid sequence (Koistinen et al., 1986).

As to the biological function of IGF-binding protein both stimulatory and inhibitory effects have been described.

Stimulatory effects of IGF-binding protein has been shown in at least two cases. Clemmons et al (1986) showed increased binding to fibroblast and smooth muscle cell surface receptors of IGF in complex with its binding protein.

Inhibitory effects of IGF-binding protein on various IGF actions in vitro, including stimulation of glucose transport by adipocytes, sulphate incorporation by chondrocytes and thymidine incorporation in fibroblasts have been described (Zapf et al., 1979; Drop et al., 1979; Ooi et al., 1986). In addition, inhibitory effects of IGF-binding proteins on growth factor mediated mitogen activity in normal cells (cartilage assay, Drop, thesis, 1983).

According to the invention, the IGF-binding protein has the following amino acid sequence:

Ala-Pro-Trp-Gln-Cys-Ala-Pro-Cys-Ser-Ala-Glu-Lys-Leu-Ala-Leu-Cys-Pro-Pro-Val-Ser-Ala-Ser-Cys-Ser-Glu-Val-Thr-Arg-Ser-Ala-Gly-Cys-Gly-Cys-Cys-Pro-Met-Cys-Ala-Leu-Pro-Leu-Gly-Ala-Ala-Cys-Gly-Val-Ala-Thr-Ala-Arg-Cys-Ala-Arg-Gly-15 Leu-Ser-Cys-Arg-Ala-Leu-Pro-Gly-Glu-Gln-Gln-Pro-Leu-His-Ala-Leu-Thr-Arg-Gly-Gln-Gly-Ala-Cys-Val-Gln-Glu-Ser-Asp-Ala-Ser-Ala-Pro-His-Ala-Ala-Glu-Ala-Gly-Ser-Pro-Glu-Ser-Pro-Glu-Ser-Thr-Glu-Ile-Thr-Glu-Glu-Glu-Leu-Leu-Asp-Asn-Phe-His-Leu-Met-Ala-Pro-Ser-Glu-Glu-Asp-His-Ser-Ile-Leu-20 Trp-Asp-Ala-Ile-Ser-Thr-Tyr-Asp-Gly-Ser-Lys-Ala-Leu-His-Val-Thr-Asn-Ile-Lys-Lys-Trp-Lys-Glu-Pro-Cys-Arg-Ile-Glu-Leu-Tyr-Arg-Val-Val-Glu-Ser-Leu-Ala-Lys-Ala-Gln-Glu-Thr-Ser-Gly-Glu-Glu-Ile-Ser-Lys-Phe-Tyr-Leu-Pro-Asn-Cys-Asn-Lys-Asn-Gly-Phe-Tyr-His-Ser-Arg-Gln-Cys-Glu-Thr-Ser-Met-25 Asp-Gly-Glu-Ala-Gly-Leu-Cys-Trp-Cys-Val-Tyr-Phe-Trp-Asn-Gly-Lys-Arg-Ile-Pro-Gly-Ser-Pro-Glu-Ile-Arg-Gly-Asp-Pro-Asn-Cys-Gln-Met-Tyr-Phe-Asn-Val-Gln-Asn

The complete nucleotide sequence of the corresponding cDNA sequence was shown to have the structure depicted in claim 5.

The protein according to the invention may be used as an effective potentiator for the functioning of

somatomedins. The effect can be mediated through the firm binding between the somatomedins and their binding proteins under physiological conditions. Such complexed somatomedins together with their binding proteins are protected against undue proteolysis, causing a significant increase of the biological half life of somatomedins. Furthermore, this IGF-binding protein or modifications hereof might function as a potent carrier of IGF to its local sites of action.

- The studies of Clemmons (1986) demonstrate the potential usefulness of this IGF-1 binding protein or modifications hereof in fertilization and in potentiation of growth of connective tissue and muscle cells in tissue repair.
- As IGF-binding protein, or modifications thereof, such
 as alpha 1 PEG, are the major secretory soluble protein
 of decidual cells of the endometirum, IBP-1 may have
 an important function in restricting trophoblast invasion
 into the endometrium during placental development.
 Furthermore, the inhibitory function of IBP-1 in cellular
 proliferation assays and the unexpected direct inhibitory
 effect of IBP-1 on the oestrogen response on certain
 cancer cells make IBP-1 or modifications hereof a potential anticancer reagent with local growth inhibitory
 effect.
- The invention is explained more in detail in the following description with reference to the drawing, in which
 - fig. 1 illustrates a restriction map and sequence strategy for human 28 kD IGF binding protein cDNA clones,
- fig. 2 shows the nucleotide and deducted amino acid sequence of human IGF binding protein where differences between the placental cDNA sequence and the liver cDNA sequence are shown in parenthesis, and

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fig. 3 represents an SDS/PAGE analysis of culture media of COS-1 cells transfected with pSV19, pSV4, pSV4Inv and untransfected COS-1 cells.

The cDNA encoding IGF-BP was obtained by screening a human placental and a human hepatoma (HEPG-2) cDNA-expression library with a polyclonal antibody to human amniotic fluid binding protein (Drop et al., 1984a).

Restriction analysis indicated that the clones isolated from the placenta library and the clones isolated from the HEPG2 library were colinear (fig. 1), supporting their candidacy as IGF binding protein clones. The composite restriction map is shown at the top of the figure 1. A putative leader sequence is shown in front of the open box representing translated regions. Each arrow shows the direction and extend of sequencing by chain termination. Four different clones are aligned. p4 and p19 originate from the placental cDNA library, while w61 and w85 originate from the HEPG2 cDNA library, (E = EcoRI, P = Pstl, B = BamHI, H = HindII, S = Sstl, X = Xbal, N = Ncol).

The complete nucleotide sequence of the cDNA insert of one of the clones isolated (pl9) was determined. The 1421 nucleotide sequence shown in fig. 2 contains a 5' untranslated region of 52 nucleotides followed by an ATG codon and an open reading frame of 776 nucleotides. The potential initiation codon is flanked by 5 sequences matching Kozak's criteria for an initiation codon (Kozak, 1986). At the 3' end the open reading frame is flanked by a translation termination codon (TGA) and a 569 nucleotides long 3' untranslated sequence.

The open reading frame in cDNA clone pl9 has a coding capacity for a protein of 259 residues also shown in

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figure 2 (by the one-letter code), with a calculated Mw of 28,172 daltons. The initiation methione is the first amino acid of a 24-residue highly hydrophobic peptide sequence (underlined), representing the sequence 5 of a putative signal sequence necessary for transfer of the nascent polypeptide sequence into the membranes of the endoplasmatic reticulim. A favourable signal peptidase cleavage site (ala-gly) occurs immediately N-terminally of the alanine residue at pos +1 (von 10 Heijne, 1987). The NH_2 -terminus of the predicted mature protein is identical to the chemically determined NH2terminus described for the IGF-binding protein isolated both from amniotic fluid (Povoa et al., 1984), and from the HEPG2 cell line (Povoa et al., 1985) and from 15 serum (Baxter et al., 1987).

Omitting the signal peptide sequence, the M_{r} of this gene product is predicted to be 2,350 daltons. The M_{r} of serum IGF binding protein is about 28,000 daltons (Baxter et al., 1987). The difference is believed to be accounted for by glycosylation of the IGF binding protein (Bohn et al., 1980; Koistinen et al., 1986).

The amino acid sequence did not disclose N-linked glycosylation sites (N-T, N-S). However, at least five potential O-linked glycosylation sites were found in the NH2-terminal of the molecule. A RGD sequence in the COOH terminal part of the IBP-l protein has been found. Such a short sequence is considered to be important for cellular attachment of matrix proteins, such as fibronectin, vitronectin and von Willebrand factor, to receptors of the integrin family.

Amino acid homology with other known proteins and peptides were determined by searching the NFBR data base, version 12.0 and 26.0. The IGF-BP protein did not show

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any significant protein homology, indicating that IGF-BP is a unique protein. In particular, comparison of amino acid sequence of IGF-I, IGF-II, Insulin, Type I IGF receptor and the type II IGF receptor amino acid sequences revealed no homologous domains. In addition, no homology was found with the reported NH₂-terminal amino acid sequence for the high molecular weight IGF binding protein (Baxter et al., 1986).

It was further possible to express IBP-1 in mammalian 10 cells. Expression vectors, pSV19, pSV4 and pSV4Inv were constructed by inserting the full length clones p4 and p19 in the expression vector pSV328. The vectors pSV4, pSV19 and pSV4Inv, in which the cDNA insert is in 3' - 5' orientation, were transfected to COS-1 cells. 15 By use of SDS/PAGE analysis cell culture media from COS-1 cells transfected with pSV19 (lane A), pSV4 (lane B), and amniotic fluid (lane C) were analysed. The IGF binding proteins were made visible by immuno staining as described for the screening of the cDNA libraries. 20 In culture media of pSV4 and pSV19 transfected COS-1 cells in which the gene is in the correct orientation a protein of 32 kD being immunologically indistinguishable from the IGF binding protein from amniotic fluid (fig. 3) was detected. In all culture media a band 25 was visible which reacted with the 35 kD SMBP antibody but which was absent in the culture medium from untransfected COS-1 cells.

Although IBP-1 successfully has been expressed in COS-1 cells the lack of N-linked glycosylation sites in the putative protein also favour expression in yeast and bacteria to increase the IBP-1 production to be used in a variety of therapeutic compositions.

The invention provides therapeutic compositions comprising IBP-1 or derivatives thereof and pharmacologically acceptable excipients. Such compositions including the IGF-binding protein or derivatives hereof according to this invention have many therapeutic uses involving the physiological functions of somatomedins.

- The IGF-binding protein of the invention may be formulated as pharmaceutical preparations comprising the IGF-binding protein of the invention together with the usual excipients. Pharmaceutical preparations according to the invention may be in the form of suspension or solutions for parentheral administration, e.g. i.v., s.c., i.m., implants, subcutaneous or interveneous administration or administration through the mucosa, e.g. oral, nasal, buccal, sublingual or rectal administration or transdermal administration.
- 15 For example in cases where somatomedins have toobe transported to specific target tissues in a way where the physiological halflife of the somatomedins has to be increased by complexing IBP-1 described in this invention to IGF-1 and IGF-2. In accordance with this 20 invention a slow release of active IGF-1 or 2 from such complexes would ascertain a constant level of somatomedins either locally or systemically dependent upon the way of administration. IBP-1 describes in this invention hereby abolishes the potent mitogenic 25 effect of the somatomedins that administrated in high dosis, i.e. intra venously, would cause unwanted local cellular proliferations in a variety of cells like fibroblasts, muscle cells and endothelial cells.
- However, the IBP-1 described by this invention administred together with IGF-1, IGF-2 and other growth factors or formulated as common preparations for topical
 use (such as PDGF, EGF, FGF, TGFalpha or TGFbetha)
 employed in therapeutical devices to be used in healing
 of wounds or in treatment of oeteoporosis and in healing

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of bones might be valuable for a steady and controlled release of the somatomedins in such therapeutical devices.

Such preparations may optionally be administred in the form of combination preparations e.g. comprising IBP-1 and IGF-1, IBP-1 and IGF-2 or IBP-1, IGF-1 and IGF-2.

In general, IBP-1 according to this invention might turn out to exhibit a potent regulatory function in the release of IGF-1 and/or IGF-2 in future treatment of injuries or other malfunctions that requires increased IGF-1 and/or IGF-1 levels.

On the other hand IBP-1 or derivatives thereof according to this invention might be useful in therapy of the proliferation of certain cancers characterized by producing somatomedins in high amounts thus inhibiting the autocrine/paracrine physiological stimulation of unwanted cellular proliferation in cancers like chondrosarcomas, fibrosarcomas, and mammacarcinomas.

Furthermore, the IBP-1 or derivatives hereof described in this invention is useful for the production of antibodies. Such mono- or polyclonal antibodies are suitable for developing immunological methods like immunohisto-chemical analysis of IBP-1 in tissues and for developing ELISA for IBP-1 quantitation. Such ELISA will prove valuable for early screening the levels of IBP-1 in patients with altered IGF-1 and 2 levels.

Pharmaceutical preparations of this invention for s.c. and i.m. administration can be prepared by mixing the following constituents: IBP-1 and derivatives thereof together with IGF-1, IGF-2 and other growth factors, an isotonic agent, a buffer, a preservative and water.

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After mixing the pH value of the preparation is, if necessary, adjusted to pH = 7.3.

Examples of preservatives: phenol and m-cresol. Examples of an isotonic agent: sodium chloride and glycerol. Example of buffer is sodium phosphate.

Pharmaceutical preparations of this invention for transmucosal administration can be prepared by mixing the following constituents: IBP-1 and derivatives thereof, together with IGF-1, IGF-2 and other growth factors, a buffer, an isotonic agent, a preservative, an absorption promotor and a vehicle e.g. water, cellulose, water-soluble cellulose alkylethers, crystalline cellulose, water-soluble polyacrylates or mixtures thereof.

Pharmaceutical preparations of this invention for transdermal administration can be prepared by mixing the
following constituents: IBP-1 and derivatives thereof
together with IGF-1, IGF-2 and other growth factors,
an isotonic agent, a preservative and a vehicle e.g.
a hydrophilic gel of water-soluble cellulose alkylethers.

This invention is further explained in the following working example describing the isolation and characterization of IBP-1.

EXAMPLE

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25 Screening the Lambda gtll Expression Library

A human placenta cDNA library in lambda gtll and a cDNA library of the human hepatoma cell line HEPG2 were screened with a polyclonal antibody to human amniotic fluid binding protein according to the procedure described by Young and Davis (Young and Davis, 1982).

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Rabbit antibody to 35 kD somatomedin binding protein SMBP isolated from human amniotic fluid was produced and purified as described by Drop et al., 1984a. The antibody was absorbed against E.coli Y1090 and lambda gtll proteins by incubating with nitrocellulose filters that had been lifted from confluent lysis plates of E.coli Y1090/lambda gtll induced with 10 mM isopropyl beta-d-thiogalacopyranoside (IPTG). The antibody was further absorbed against human serum albumin immobilized on nitrocellulose filters. Approximately 4×10^5 clones of the placental library were screened and about 0.5 x 10⁵ of the HPEG2 library. 3-5 x 10⁴ plague forming units per 150 mm Petri dish were plated on a lawn of Y1090 bacteria and incubated. After 2 hr incubation the plates were covered with nitrocellulose filters (Millipore HATF) that had been saturated with 10 mM IPTG. The plates were incubated at 37°C for 2-2.5 hrs. The filters were removed, washed with Tris-buffered saline (TBS; 10 mM Tris/HCl, pH 7.5/150 mM NaCl) at room temperature and incubated with 3% BSA in TBS for 30 min. at room temperature. Partly purified rabbit polyclonal 35 kD SMBP antibody diluted 1:125 was added to 3% BSA in TBS plus 0.02% azide, and the filters were incubated overnight at 4°C. The filters were washed and incubated for 60 min. at room temperature with horse-radish peroxydase conjugated goat anti-rabbit IgG (Tago) diluted 1:200 in 3% BSA in TBS. The filters were washed and stained with amidophenyl and napthol AS-MX phosphate in 0.2 M Tris/HCl, pH 9.2, 10 mM MgCl, at room temperature.

Positive phages were isolated and DNA was isolated by standard methods (Maniatis et al., 1982). About 33 plaques strongly cross-reacting with the polyclonal antibody were identified in the placenta and HEPG2 cDNA library. Following re-screening inserts varying in size between 0.9-1.5 Kb were isolated and subcloned

in the vector PTZ19 from Pharmacia. All isolated clones showed cross-hybridization in a Southern blot except one clone from the placenta library and the 5 weakly hybridizing clones from the HEPG2 library.

DNA was digested with various restriction endonucleases (BRL, NEN, Boehringer) according to the suppliers directions, electrophoresed in 0.8% agarose, and transferred to nitrocellulose filters according to the method of southern (Southern, 1975). mRNA was denaturated with dimethylsulfozide (DMSO) and glyoxal, subjected to electrophoresis in 1% agarose and transferred to nitrocellulose filters (Millipore HFTF).

Restriction fragments were subcloned in the vectors PTZ18 or PTZ19 (Pharmacia) and sequenced according to the chain termination method (Sanger et al., 1977). In regions which lacked convenient restriction sites, appropriate clones were generated by Bal 31 nuclease digestion.

Transfection of COS-1 Cells

20 The full length cDNA clones p4 and p19 were subcloned in the EcoRl site of pSV328, which expressed cloned inserts using the simian virus 40 (SV40) early promotor (Van Heuvel et al., 1986). A DEAE-dextran procedure (McCuthchan & Pagano, 1986) followed by treatment with 100 uM chloroquine in Dulbecco's MEM (DMEM) for 4 hrs 25 was used to transfect COS-1 cells (Gluzman, 1981). After this treatment the cells were grown 24 hrs with DMEM plus 5% foetal calf serum. Medium was removed after 72 hrs, and the cells were washed extensively with DMEM and incubated for 72 hrs with DMEM without 30 serum. Production of 32 kD binding protein in culture media was determined using 35 kD SMBP antibody.

Purification of IBP-1

Proteins from amniotic fluid or from conditioned media were precipitated with ammonium sulphate at a final concentration of 35%. Following centrifugation the 5 supernatant was brought to 50% ammonium sulphate. The pellet was dissolved in 45% ammonium sulphate and the final pellet was dissolved in 50 mM Tris HCl, pH 7.5 for further purification and characterization. The dissolved ammonium sulphate precipitate was further 10 purified by reverse phase chromatography on C18. Following washings with 50 mM Tris-HCl, pH 7.5 and Tris HCl, pH 7.5 in 50% methanol the pure IBP-1 was eluted from the column with Tris-HCl, pH 7.5 in 65% methanol. The IBP-1 was precipitated overnight in a 7% Trichloro-15 acetic-acid solution. The precipitate was dissolved in 20 mM Tris-HCl, pH 7.5, lyophilized and storred at ÷ 20°C.

Characterization of purified IBP-1

The biological effects of purified samples if IBP-l were tested in in vitro mitogenicity tests employing ³H-thymidin incorporation into a MCF-7 mamma carcinoma cell line. The stimulatory effect of both IGF-l and

IGF-2 on cell proliferation was inhibited by IBP-1 in a dose dependent way. The effect of IBP-1 on IGF-2 dependent cell proliferation was more pronounced than that of the effect of IBF-1.

Furthermore, in an assay employing | 35S|-Methionin incorporation into cartillage the stimulatory effect of IGF-1 and IGF-2 was abolished by IBP in the low ng range.

Examples of pharmaceutical preparations:

10 EXAMPLE 1

25 ng IBP-1

10 ng IGF-1

0.7% NaCl

1/75M sodiumphosphate

15 water ad 1 ml

The calculated amounts of IBP-1 and IGF-1 were dissolved and diluted in phosphate buffer containing NaCl. The pH was adjusted to 7.3-7.4.

EXAMPLE 2

20 25 ng IBP-1

10 ng IGF-2

1.6% glycerin

1/75M sodiumphosphate

0.01% benzalconiumchlorid

25 0.05% sodiumedetat

water ad I ml

The calculated amounts of IBP-1 and IGF-2 were dissolved and diluted in phosphate buffer containing glycerin, benzalconiumchlorid and sodiumedetat. The pH NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENT AND/OR AMINO ACID SEQUENT DISCLOSURES

The nucleotide and/or amino acid sequence disclosure contained in this application does not
comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the
following reason(s):

	1. This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to these regulations, published at 1114 OG 29, May 15, 1990 and at 55 FR 18230, May 1, 1990.
	2. This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c).
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	4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing."
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	6. The paper copy of the "Sequence Listing" is not the same as the computer readable from of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).
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was adjusted to 7.4

EXAMPLE 3

- 25 ng IBP-1
- 10 ng IGF-1
- 5 10 ng IGF-2
 - 5% hydroxyethylcellulose
 - 0.9% benzylalcohol
 - 1/75M phosphate buffer

water ad 1 ml

The gel is prepared by mixing hydroxyethylcellulose with the waterphase containing IBP-1, IGF-1 and IGF-2.

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Patent Claims:

1. An IGF binding protein, comprising the following amino acid sequence:

A-P-W-Q-C-A-P-C-S-A-E-K-L-A-L-C-P-P-V-S-A-S-C-S-E-V-T-R-S-A
G-C-G-C-C-P-M-C-A-L-P-L-G-A-A-C-G-V-A-T-A-R-C-A-R-G-L-S-C-R-A-L-P-G-E-Q-Q-P-L-H-A-L-T-R-G-Q-G-A-C-V-Q-E-S-D-A-S-A-P-H-A-A-E-A-G-S-P-E-S-P-E-S-T-E-I-T-E-E-E-L-L-D-N-F-H-L-M-A-P-S-E-E-D-H-S-I-L-W-D-A-I-S-T-Y-D-G-S-K-A-L-H-V-T-N-I-K-K-W-E-P-C-R-I-E-L-Y-R-V-V-E-S-L-A-K-A-Q-E-T-S-G-E-E-I-S-K-F-Y-L-P-N-C-

10 N-K-N-G-F-Y-H-S-R-Q-C-E-T-S-M-D-G-E-A-G-L-C-W-C-V-Y-P-W-N-G-K-R-I-P-G-S-P-E-I-R-G-D-P-N-C-Q-M-Y-F-N-V-Q-N-

or an equivalent modification thereof.

- 2. A glycosylated modification of the protein according to claim 1.
- 3. An IGF binding protein according to claim 2, in which one or more hydroxy groups are glycosylated.
 - 4. A DNA sequence, coding for the IGF binding protein as defined in claim 1.
- 5. A cDNA sequence according to claim 4 in which the coding strand includes the following structure:

				•	
1	GGGCGGCAC	AGCCAGAGAG	CATCGGCCCC	TETCTECTEC	TCGCGCCTGG
51	AGATGITCAGA	GGTCCCCGTT	GCTCGCGTCT	GECTESTACT	GCTCCTGCTG
101	ACTGTCCAGG	TCGGCGTGAC	AGCCGGCGCT	CCGTGGCAGT	GCGCGCCCTG
151	CTCCGCCGAG	AAGCTCGCGC	TCTGCCCGCC	GGTGTCCGCC	TCGTGCTCGG
201	AGGTCACCCG	GTCCGCCGGC	TGCGGCTGTT	GCCCGATGTG	CGCCCTGCCT
251	CTGGGCGCCG	CGTGCGGCGT	GGCGACTGCA	CGCTGCGCCC	GGGGACTCAG
301	TTGCCGCGCG	CTGCCGGGGG	AGCAGCAACC	TCTGCACGCC	CTCACCCGCG
351	GCCAAGGCGC	CTGCGTGCAG	GAGTCTGACG	CCTCCGCTCC	CCATGCTGCA
401	GAGGCAGGGA	GCCCTGAAAG	CCCAGAGAGC	ACGGAGATAA	CTGAGGAGGA
451	GCTCCTGGAT	AATTTCCATC	TEATEGCCCC	TTCTGAAGAG	GATCATTCCA
501	TCCTTTEGGA	CGCCATCAGT	ACCTATGATG	GCTCGAAGGC	TCTCCATGTC
551	ACCAACATCA	AAAAATGGAA	GGAGCCCTGC-	CGAATAGAAC	TCTACAGAGT
601	CGTAGAGAGT	TTAGCCAAGG	CACAGGAGAC	ATCAGGAGAA	GAAATTTCCA
451	AATTTTACCT	GCCAAACTGC	AACAAGAATG	GATTTTATCA	CAGCAGACAG
701	TGTGAGACAT	CCATGGATEG	AGAGGCGGGA	CTCTGCTGGT	GCGTCTACCC
751	TTGGAATGGG	AAGAGGATCC	CTGGGTCTCC	AGAGATCAGG	GGAGACCCCA
801	ACTGCCAGAT	GTATTTTAAT	GTACAAAACT	GAACCAGAT	GAAATAATGT
851	TCTGTCACGT	GAAATATTTA	AGTATATAGT	ATATTTATAC	TCTAGAACAT
901	GCACATTTAT	ATATATGTAT	ATGTATATAT	ATATAGTAAC	TACTTCTTAT
951	ACTCCATACA	TAACTTGATA	TAGAAAGCTG	TTTATTTATT	CACTETAAGT
1001	TTATTTTTC	TACACAGTAA	AAACTTGTAC	TATGTTAATA	ACTTGTCCTA
1051	TGTCAATTTG	TATATCATGA	AACACTTCTC	ATCATATTGT	ATGTAAGTAA
1101	TTGCATTTCT	GCTCTTCCAA	AGCTCCTGCG	TCTGTTTTTA	AAGAGCATGG
1151	AAAAATACTG	CCTAGAAAAT	GCAAAATGAA	ATAAGAGAGA	GTAGTTTTTC
1201	AGCTAGTTTG	AAGGAGGACG	GTTAACTTGT	ATATTCCACC	ATTCACATTT
1251	GATGTACATG	TGTAGGGAAA	GTTAAAAGTG	TTGATTACAT	AATCAAAGCT
1301	ACCTGTGGTG	ATAGTTGCCA	CCTGTTAAAA	TGTACACTGG	ATATGTTGTT
1351	AAACACGTGT	CTATAATGGA	AACATTTACA	ATAAATATTC	TGCATGGAAA
1401	44444444	AAAAAAAAA	AAAAA		

- 6. An expression vector, containing the DNA sequence defined in claim 4 or 5.
- 7. The expression vector pSV19.
- 8. The expression vector pSV4.
- 9. A cell line or a microorganism, comprising an expression vector in accordance with claims 6, 7 or 8.
 - 10. A pharmaceutical preparation, comprising the binding protein defined in any of the claims 1-3.

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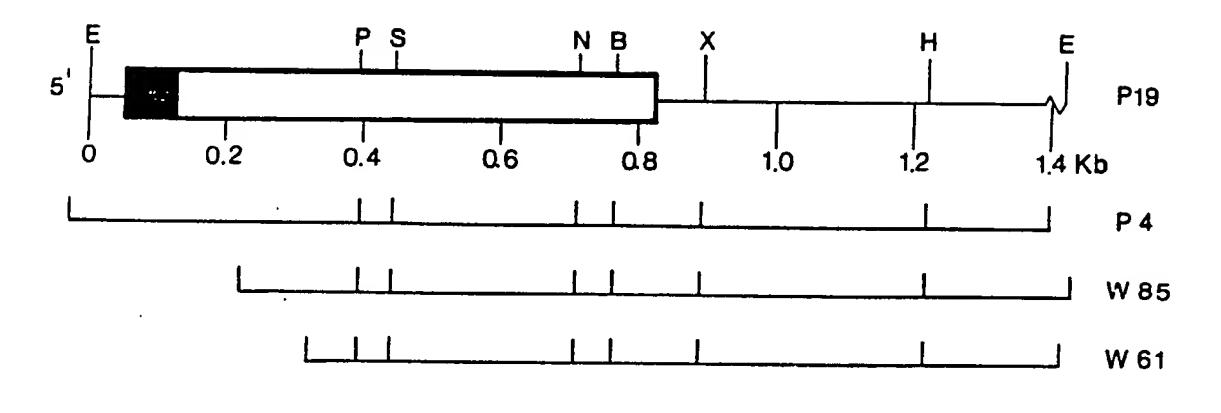


FIG.1

					-25				
GGGCGGGCACA 10	GCCAGAGAGCA 20	TCGGCCCCTGT(30	CTGCTGCTC 40	GCGCCTGGAG 50	M S E ETGTCAGAGG 60	TCCCCGTTGC 70	TCGCGTCTGC 80	GCTGGTACT 90	, +1;
L L L GCTCCTGAG	T V O V CTGTCCAGGTC 110	G V T A GGCGTGACAGCO 120	+1 <u>G</u> A P GGCGCTCC 130	W Q C GTGGCAGTGC 140		S A E K CCGCCGAGAAG 160	L A L GCTCGCGCTC 170	C P P TGCCCGCC 180	
V S A S GGTGTCCGCCTC 190	S C S E CGTGCTCGGAGO 200	V T R S STCACCCGGTCC 210	A G C GCCGGCTG(220	G C C CGGCTGTTGC 230	P M C CCGATGTGCGC	A L P L CCCTGCCTCTC 250	G A A GGGCGCCGCG 260	C G V TGCGGCGT 270	48
A T A R GGCGACTGCACG 280	R C A R SCTGCGCCCGGG 290	G L S C GACTCAGTTGC 300	R A L CGCGCGCTG 310	P G E CCGGGGGAG 320	Q Q P I CAGCAACCTCT 330	L H A L rgcacgcctto 340	T R G CACCCGCGGC 350	Q G A CAAGGCGC 360	78
C V Q E CTGCGTGCAGGA 370	S D A GTCTGACGCCT 380	S A P H CCGCTCCCCAT 390	A A E GCTGCAGAG 400	A G S GCAGGGAGC 410	PESF CCTGAAAGCCC 420	E S T CAGAGAGCACG 430	E I T GAGATAACT 440	E E E Gaggagga 450	10
L L D N GCTCCTGGATAA 460	F H L TTTCCATCTGA 470	M A P S TGGCCCCTTCT(480	E E D Gaagaggat 490	H S I CATTCCATCO 500	L W D A ETTTGGGACGC 510	I S T CATCAGTACC 520	Y D G TATGATGGC 530	S K A TCGAAGGC 540	138
L H V T TCTCCATGTCAC 550	N I K CAACATCAAAA (C)	K W K E Aatggaaggagg 570	P C R CCTGCCGA 580	I E L Atagaactci 590	Y R V V ACAGAGTCGT 600	E S L AGAGAGTTTA 610	A K A GCCAAGGCA 620	Q E T CAGGAGAC 630	168
S G E E Atcaggagaaga 640	I S K 1 AATTTCCAAAT 650	F Y L P TTTACCTGCCAA 660	N C N ACTGCAAC 670	K N G AAGAATGGAT 680	F Y H S TTTATCACAG 690	R Q C CAGACAGTGT 700	E T S Gagacatcc 710	M D G Atggatgg 720	198
E A G L Agaggcgggacto 730	C W C T CTGCTGGTGCGT 740	V Y P W Ictaccttgga 750	N G K ATGGGAAG 760	•	G S P E GGTCTCCAGA 780		D P N GACCCCAAC 800	C Q M IGCCAGAT 810	228
Y F N V GTATTTTAATGTA 820	Q N ACAAAACTGAAA 830	ACCAGATGAAAT 840	AATGTTCTC 850	STCACGTGAA 860	ATHERRAGT. 870	ATATAGTATE: 880	MEDITACTC1 890	TAGAACAT 900	234
GCACAGRETATA 910	TATGTATATG7 920	ATATATATATA 930	GTAACTACT 940	TCTTATACT 950	CCATACATAA 960	CTTGATATAGA 970	AAGCTGTTT 980	PERFECT 990	
CACTGTAAGTTTA 1000	TTTTTTTCTACA 1010	CAGTAAAAACT 1020 j	TGTACTATG 1030	TTAATAACT 1040	IGTCCTATGTO 1050	CAATTTGTATA 1060	ATCATGAAAC 1070	ACTTCTC (T)1080	•
ATCATATTGTATG 1090	TAAGTAATTGC 1100	ATTTCTGCTCT' 1110	ICCAAAGCT (G)	CCTGCGTCT(1130	TTTTTAAAGA 1140 (C		ATACTGCCT 1160	AGAAAAT 1170	
CAAAATGAAATA 1180	AGAGAGAGTAG 1190	TTTTTCAGCTAG	STTTGAAGG 1210	AGGACGGTT/ 1220	ACTTGTATAT 1230	TCCACCATTC 1240	ACATTTGAT	GTACATG 1260	
GTAGGGAAAGTT 1270	AAAAGTGTTGA 1280	TTACATAATCAA 1290	AGCTACCT	GTGGTGATAC 1310	TTGCCACCTG 1320	TTAAAATGTA 1330	CACTGGATA' 1340	1350	
AACACGTGTCTA 1360	TAATGGAAAC#	TETROAATAAA) 1380	TATTCTGCA' 1390	TGGAAAAAA 1400	.AAAAAAAAAA 1410	AAAAAAAAA 1420	A		

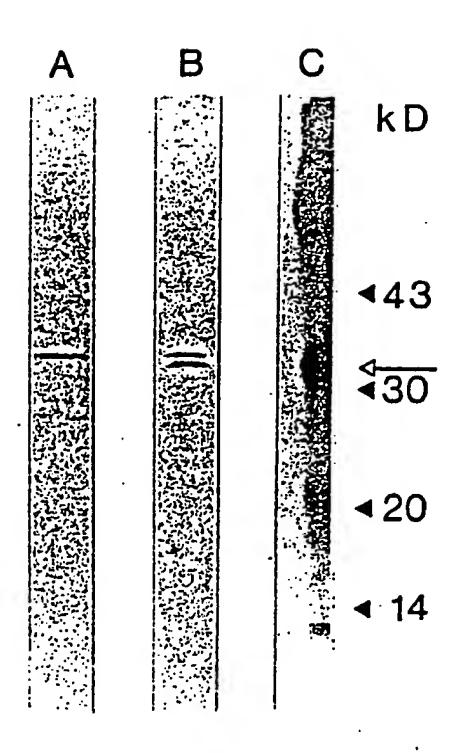


FIG. 3

INTERNATIONAL SEARCH REPORT

International Application No PCT/NL 89/00008

I. CLASS	SIFICATION OF SUBJECT MATTER (it several class	ification sympols apply, indicate all) 5	
According	to International Patent Classification (IPC) or to both Nat		
IPC4:	C 07 K 13/00, C 12 N 15/00 //A 61 K 35/50	O, C 12 P 21/02, A	61 K 37/02,
II. FIELD	S SEARCHED		
	Minlmum Docume	ntation Searched 7	
Classificati	on System	Classification Symbols	
IPC ⁴	C 07 K; C 12 N; C 1	L2 P; A 61 K	
	Documentation Searched other to the Extent that such Documents	than Minimum Documentation are included in the Fields Searched	
III. DOCL	Citation of Document, 19 with Indication, where app	propriate, of the relevant passages 12	Relevant to Claim No. 13
X	EP, A, 0141326 (BEHRINGWE 15 May 1985, see the	ERKE)	1-3,10
X	Biochemical and Biophysic Communications, vol. 23 November 1986, Aca (US) M.L. Huhtala et active domain in soma protein", pages 263-2 see the whole article	141, no. 1, demic Press, Inc. al.: "Biologically tomedin-binding 270	1-3,10
P,X	Molecular Endocrinology, May 1988 The Endrocrine Societ Y.L. Lee et al.: "Ins growth factor (IGF) b complementary deoxyri	y ulin-like inding protein bonucleic acid	1-10
	from human HEP G2 hep predicted protein seq IGF binding domain di of the IGF-I and IGF-pages 404-411 see the whole article	uence suggests an fferent from those II receptors"	
"A" doct cont filing "L" doct which citat "O" doct othe "P" doct later	i categories of cited documents: 10 ument defining the general state of the art which is not sidered to be of particular relevance ier document but published on or after the International grate ument which may throw doubts on priority claim(s) or the cited to establish the publication date of another iron or other special reason (as specified) ument referring to an oral disclosure, use, exhibition or ir means ument published prior to the international filling date but than the priority date claimed	"T" fater document published after the or priority date and not in conflic cited to understand the principle invention "X" document of particular relevance cannot be considered novel of involve an inventive step "Y" document of particular relevance cannot be considered to involve a document is combined with one of ments, such combination being of in the art. "4" document member of the same priority of	or theory underlying the e; the claimed invention cannot be considered to e; the claimed invention inventive step when the or more other such docu- bytous to a person skilled etent family
Date of the	Actual Completion of the International Search	4 1. 07. 89	Hen waken
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	Citation of Document, with indication, where appropriate, of the relevant passages	Relevant to Claim No
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P,X	Embo Journal, vol. 7, no. 8, 1988 IRL Press Ltd. Oxford (GB) A. Brinkman et al.: "Isolation and characterization of a cDNA encoding the low molecular weight insulin-like growth factor binding protein (IBP-1) pages 2417-2423	1-10
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ANNEX TO THE INTERNATIONAL SEARCH REPORT ON INTERNATIONAL PATENT APPLICATION NO.

NL 8900008 SA 27516

This annex lists the patent family members relating to the patent documents cited in the above-mentioned international search report. The members are as contained in the European Patent Office EDP file on 05/07/89

The European Patent Office is in no way liable for these particulars which are merely given for the purpose of information.

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